



SEQUENCE LISTING

<110> YEH, EDWARD T.H.

<120> USES FOR A NOVEL CELL-DEATH-PROTECTING PROTEIN

<130> UTSH:248US

<140> 09/484,964

<141> 2000-01-18

<150> 08/964,162

<151> 1997-11-04

<150> 60/030,302

<151> 1996-11-05

<160> 18

<170> PatentIn Ver. 2.0

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<211> 1465

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (88)..(390)

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Met Ser Asp Gln Glu Ala Lys Pro Ser

1

5

act gag gac ttg ggg gat aag aag caa ggt gaa tat att aaa ctc aaa 162  
Thr Glu Asp Leu Gly Asp Lys Gln Gly Glu Tyr Ile Lys Leu Lys  
10 15 20 25

gtc att gga cag gat agc agt gag att cac ttc aaa gtg aaa atg aca 210  
Val Ile Gly Gln Asp Ser Ser Glu Ile His Phe Lys Val Lys Met Thr  
30 35 40

aca cat ctc aag aaa ctc aaa gaa tca tac tgt caa aga cag ggt gtt 258  
Thr His Leu Lys Leu Lys Glu Ser Tyr Cys Gln Arg Gln Gly Val  
45 50 55

cca atg aat tca ctc agg ttt ctc ttt gag ggt cag aga att gct gat 306  
Pro Met Asn Ser Leu Arg Phe Leu Phe Glu Gly Gln Arg Ile Ala Asp  
60 65 70

aat cat act cca aaa gaa ctg gga atg gag gaa gaa gat gtg att gaa 354  
Asn His Thr Pro Lys Glu Leu Gly Met Glu Glu Asp Val Ile Glu  
75 80 85

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Val Tyr Gln Glu Gln Thr Gly Gly His Ser Thr Val  
90 95 100

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gaattctagt gctcattatt cattattgtt tgttttcatt gtgctgattt ttgggtatca 580  
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<211> 101  
<212> PRT  
<213> Homo sapiens

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20 25 30  
Glu Ile His Phe Lys Val Lys Met Thr Thr His Leu Lys Lys Leu Lys  
35 40 45  
Glu Ser Tyr Cys Gln Arg Gln Gly Val Pro Met Asn Ser Leu Arg Phe  
50 55 60  
Leu Phe Glu Gly Gln Arg Ile Ala Asp Asn His Thr Pro Lys Glu Leu  
65 70 75 80  
Gly Met Glu Glu Glu Asp Val Ile Glu Val Tyr Gln Glu Gln Thr Gly  
85 90 95  
Gly His Ser Thr Val  
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<223> N = A, C, G or T  
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 cagtttaga ttaaggaggca tacaccactt agtaaactaa taaaagccta ttgtgaacga 240  
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 <212> PRT  
 <213> Homo sapiens

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 20 25 30  
 Lys Ile Lys Arg His Thr Pro Leu Ser Lys Leu Met Lys Ala Tyr Cys  
 35 40 45  
 Glu Arg Gln Gly Leu Ser Met Arg Gln Ile Arg Phe Arg Phe Asp Gly  
 50 55 60  
 Gln Pro Ile Asn Glu Thr Asp Thr Pro Ala Gln Leu Glu Met Glu Asp  
 65 70 75 80  
 Glu Asp Thr Ile Asp Val Phe Gln Gln Gln Thr Gly Gly Val Tyr  
 85 90 95

<210> 5  
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 <212> DNA  
 <213> Homo sapiens  
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 <221> modified\_base  
 <222> (19)  
 <223> N = A, C, G or T

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cacaaccaaa attggccctt ttctccctt taatattgaa gaaattccca catttctcat 1200  
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aagagctact tccaaatgtg gttacaaatg aaccatggaa atgatgactt catgttcttc 1380  
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<212> PRT  
<213> Homo sapiens

<400> 6  
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20 25 30  
Ile Lys Arg His Thr Ser Leu Ser Lys Leu Met Lys Ala Tyr Cys Glu  
35 40 45  
Arg Gln Gly Leu Ser Met Arg Gln Ile Arg Phe Arg Phe Asp Gly Gln  
50 55 60  
Pro Ile Asn Glu Thr Asp Thr Pro Ala Gln Leu Arg Met Glu Asp Glu  
65 70 75 80  
Asp Thr Ile Asp Val Phe Gln Gln Gln Thr Gly Gly Val Pro Glu Ser  
85 90 95  
Ser Leu Ala Gly His Ser Phe  
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<210> 7  
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<212> PRT  
<213> Artificial Sequence  
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<223> Description of Artificial Sequence: Synthetic Peptide

<400> 7  
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<210> 8  
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<212> DNA  
<213> Homo sapiens

<400> 8  
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<210> 9  
<211> 30  
<212> DNA  
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<400> 9  
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<210> 10  
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<213> Artificial Sequence  
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<223> Description of Artificial Sequence: Synthetic Peptide

<400> 10  
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<210> 11  
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<213> Influenza virus

<400> 11  
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<210> 12  
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<212> PRT  
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<400> 12  
His Ser Thr Val  
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<210> 13  
<211> 101  
<212> PRT  
<213> Saccharomyces cerevisiae

<400> 13  
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20 25 30  
Ser Glu Ile Phe Phe Lys Ile Lys Lys Thr Thr Pro Leu Arg Arg Leu  
35 40 45  
Met Glu Ala Phe Ala Lys Arg Gln Gly Lys Glu Met Asp Ser Leu Arg  
50 55 60  
Phe Leu Tyr Asp Gly Ile Arg Ile Gln Ala Asp Gln Thr Pro Glu Asp  
65 70 75 80  
Leu Asp Met Glu Asp Asn Asp Ile Ile Glu Ala His Arg Glu Gln Ile  
85 90 95

Gly Gly Ala Thr Tyr  
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<210> 14  
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<212> PRT  
<213> Homo sapiens

<400> 14  
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20 25 30  
Lys Glu Gly Ile Pro Pro Asp Gln Gln Arg Leu Ile Phe Ala Gly Lys  
35 40 45  
Gln Leu Glu Asp Gly Arg Thr Leu Ser Asp Tyr Asn Ile Gln Lys Glu  
50 55 60  
Ser Thr Leu His Leu Val Leu Arg Leu Arg Gly Gly Gly Leu Arg  
65 70 75 80

<210> 15  
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<212> PRT  
<213> Homo sapiens

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20 25 30  
Lys Glu Gly Ile Pro Pro Gln Gln Arg Leu Ile Tyr Ser Gly Lys  
35 40 45  
Gln Met Asn Asp Glu Lys Thr Ala Ala Asp Tyr Lys Ile Leu Gly Gly  
50 55 60  
Ser Val Leu His Leu Val Leu Ala Leu Arg Gly Gly  
65 70 75

<210> 16  
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<212> PRT  
<213> Homo sapiens

<400> 16  
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<210> 17  
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<212> DNA  
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<220>

<221> CDS  
<222> (136) .. (438)

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gaagccacccg tcatac atg tct gac cag gag gca aaa cct tca act gag gac 171  
Met Ser Asp Gln Glu Ala Lys Pro Ser Thr Glu Asp  
1 5 10  
ttg ggg gat aag aag caa ggt gaa tat att aaa ctc aaa gtc att gga 219  
Leu Gly Asp Lys Lys Gln Gly Glu Tyr Ile Lys Leu Lys Val Ile Gly  
15 20 25  
cag gat agc agt gag att cac ttc aaa gtg aaa atg aca aca cat ctc 267  
Gln Asp Ser Ser Glu Ile His Phe Lys Val Lys Met Thr Thr His Leu  
30 35 40  
aag aaa ctc aaa gaa tca tac tgt caa aga cag ggt gtt cca atg aat 315  
Lys Lys Leu Lys Glu Ser Tyr Cys Gln Arg Gln Gly Val Pro Met Asn  
45 50 55 60  
tca ctc agg ttt ctc ttt gag ggt cag aga att gct gat aat cat act 363  
Ser Leu Arg Phe Leu Phe Glu Gly Gln Arg Ile Ala Asp Asn His Thr  
65 70 75  
cca aaa gaa ctg gga atg gag gaa gaa gat gtg att gaa gtt tat cag 411  
Pro Lys Glu Leu Gly Met Glu Glu Asp Val Ile Glu Val Tyr Gln  
80 85 90  
gaa caa acg ggg ggt cat tca aca gtt tagatattct ttttattttt 458  
Glu Gln Thr Gly Gly His Ser Thr Val  
95 100  
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<210> 18

<211> 101

<212> PRT

<213> Homo sapiens

<400> 18

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Lys Gln Gly Glu Tyr Ile Lys Leu Lys Val Ile Gly Gln Asp Ser Ser  
20 25 30

Glu Ile His Phe Lys Val Lys Met Thr Thr His Leu Lys Lys Leu Lys  
35 40 45

Glu Ser Tyr Cys Gln Arg Gln Gly Val Pro Met Asn Ser Leu Arg Phe  
50 55 60

Leu Phe Glu Gly Gln Arg Ile Ala Asp Asn His Thr Pro Lys Glu Leu  
65 70 75 80

Gly Met Glu Glu Glu Asp Val Ile Glu Val Tyr Gln Glu Gln Thr Gly  
85 90 95

Gly His Ser Thr Val  
100